

SPECIFICATION AMENDMENTS

Please amend the third full paragraph (lines 20 - 27) on page 9 as follows:

The 16S rDNA of #Probio-16 has the nucleotide sequence listed in SEQ. ID. NO. 1. Through a molecular systematic analysis based on 16S rDNA sequences, #Probio-16 was found to have 99.5% homology with the type strain of *Lactobacillus reuteri*, showing the highest phylogenetic relatedness among the strains tested. Based on these results, #Probio-16 was identified as a member of *Lactobacillus reuteri* strain and deposited at the Korean Culture Center of Microorganisms located at 361 – 221, Yurim B/D, Hongje-1-dong, Seodaemun-gu, Seoul, 120-091, Republic of Korea, under accession No. KCCM 10214 on Oct. 2, 2000 in accordance with the terms and provisions of the Budapest Treaty relating to deposit of microorganisms.